

OIPE

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/904,568

DATE: 07/26/2001 TIME: 18:43:12

Input Set : A:\1272591.app
Output Set: N:\CRF3\07262001\1904568.raw

### **ENTERED**

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3 <110> APPLICANT: TREMBLAY, Johanne
              HAMET, Pavel
              LEWANCZUK, Richard
      8 <120> TITLE OF INVENTION: A Novel Hypertension Related Calcium Regulated Gene
              (HCaRG)
     11 <130> FILE REFERENCE: 12725.9
     13 <140> CURRENT APPLICATION NUMBER: US/09/904,568
C--> 14 <141> CURRENT FILING DATE: 2001-07-16
     16 <160> NUMBER OF SEQ ID NOS: 4
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      34 gggaagaaag gratg tot got tig ggg got goa got coa tac tig cac cat
                      Met Ser Ala Leu Gly Ala Ala Ala Pro Tyr Leu His His
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      38 ccc gct gac agt cac agt ggc cgg gtc agt ttc ctg ggt tcc cag ccc
                                                                            218
       39 Pro Ala Asp Ser His Ser Gly Arg Val Ser Phe Leu Gly Ser Gln Pro
       42 tot coa gaa gtg acg gcc gtg gct cag ctc ttg aag gac tta gac agg
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       43 Ser Pro Glu Val Thr Ala Val Ala Gln Leu Leu Lys Asp Leu Asp Arg
       46 agc acc ttc aga aag ttg ttg aaa ctt gta gtc ggg gcc ctg cat ggg
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       47 Ser Thr Phe Arg Lys Leu Leu Lys Leu Val Val Gly Ala Leu His Gly
       50 aaa gac tgc aga gaa gct gtg gag caa ctt ggt gcc agc gcc aac ctg
                                                                             362
       51 Lys Asp Cys Arg Glu Ala Val Glu Gln Leu Gly Ala Ser Ala Asn Leu
       54 tca gaa gag cgt ctg gcc gtc ctg ctg gcg ggc aca cac acc ctg ctc
       55 Ser Glu Glu Arg Leu Ala Val Leu Leu Ala Gly Thr His Thr Leu Leu
       58 cag cag gct ctc cgg ctg ccc cct gct agt cta aag cca gat gcc ttc
       59 Gln Gln Ala Leu Arg Leu Pro Pro Ala Ser Leu Lys Pro Asp Ala Phe
        62 cag gaa gag ctc cag gaa ctt ggc att cct cag gat cta att gga gat
        63 Gln Glu Glu Leu Gln Glu Leu Gly Ile Pro Gln Asp Leu Ile Gly Asp
        66 ttg gcc agt ttg gca ttt ggg agt caa cgc cct ctt ctc gac tct gta
        67 Leu Ala Ser Leu Ala Phe Gly Ser Gln Arg Pro Leu Leu Asp Ser Val
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71 Ala Gin Gin Gin Giy Ser Sor	155
70 140	- too can tee etg caa
74 cgg gtg gac gtg gcc atc tca acc agc gct ce 75 Arg Val Asp Val Ala Ile Ser Thr Ser Ala G 165	n Ser Arg Ser Leu Gln
75 Arg Val Asp Val Ald IIe Bel III	170
76 100	t aga tot god cac cyc
78 ccg agt gtt ctc atg cag ctg aag ctc aca g 79 Pro Ser Val Leu Met Gln Leu Lys Leu Thr A	- Cly Ser Ala His Arg
76 CCg dgc 300	105
/9 PIO Sel (42 = 180	tag agt gta gcc 746
80 175 82 ttc gag gtg ccc ata gcc aaa ttt cag gag c	tg egg tac age god Ala
82 ttc gag gtg ccc ata gcc aaa ttt cag gag c 83 Phe Glu Val Pro Ile Ala Lys Phe Gln Glu L	eu Arg Tyl Scr Van
83 Phe Giu vai 110 == 195	00
84 190 195 2 86 ttg gtc ctt aag gaa atg gca gaa ctg gag a	ag aag tgt gag ege aan
86 ttg gtc ctt aag gag atg gca gaa ctg gag a 87 Leu Val Leu Lys Glu Met Ala Glu Leu Glu I 215	ys Lys Cys Giu Aig Dio
87 Leu Val Leu Lys Glu lies 1215	843
88 210 90 ctg cag gac tgactgaacc ctggtactgt gggtgct	gaa getggtacea
90 ctg cag gac tgactgaacc ctggcassy 333	003
93 gaacacagee ecceaetggt gatgageeca addititigt	tt tatgagtcaa cagctgggca 963
95 attttaagtg aaaagacage gggactttca ggttttgi 95 gggtggcaca gtttataatc tcagcccttg gaagtctc 97 gggtggcaca gtttataatc gtgaggctca gtgtcga	att aaagaggtaa agcaactatt 1083
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105 <2115 LENGTH: 224 106 <2125 TYPE: PRT 107 <213> ORGANISM: Rattus rattus 109 <400> SEQUENCE: 2 110 Met Ser Ala Leu Gly Ala Ala Ala Pro Tyr 111 1 1 5 13 Ser His Ser Gly Arg Val Ser Phe Leu Gly 114 20 115 Val Thr Ala Val Ala Gln Leu Leu Lys Asg 116 Val Thr Ala Val Ala Gln Leu Leu Lys Asg 117 35 119 Arg Lys Leu Leu Lys Leu Val Val Gly Ala 120 50 122 Arg Glu Ala Val Glu Gln Leu Gly Ala Se 123 65 125 Arg Leu Ala Val Leu Eeu Ala Gly Thr Hi 126 85 127 61 100 127 100 131 Leu Gln Glu Leu Gly Ile Pro Gln Asg Leu 132 115 133 130 135 135 130 135 137 Gln Gly Ser Ser Leu Pro His Val Ser Ty 137 Gln Gly Ser Ser Leu Pro His Val Ser Ty	Ser Gln Pro Ser Pro Glu 30  Leu Asp Arg Ser Thr Phe 45  Leu His Gly Lys Asp Cys 60  r Ala Asn Leu Ser Glu Glu 75  80  s Thr Leu Leu Gln Gln Ala 95  O Asp Ala Phe Gln Glu Glu 110  u Ile Gly Asp Leu Ala Ser 125  u Asp Ser Val Ala Gln Gln 140  or Phe Arg Trp Arg Val Asp 160
105 <2115 LENGTH: 224 106 <2125 TYPE: PRT 107 <213> ORGANISM: Rattus rattus 109 <400> SEQUENCE: 2 110 Met Ser Ala Leu Gly Ala Ala Ala Pro Tyr 111 1 1 5 13 Ser His Ser Gly Arg Val Ser Phe Leu Gly 114 20 115 Val Thr Ala Val Ala Gln Leu Leu Lys Asg 116 Val Thr Ala Val Ala Gln Leu Leu Lys Asg 117 35 119 Arg Lys Leu Leu Lys Leu Val Val Gly Ala 120 50 122 Arg Glu Ala Val Glu Gln Leu Gly Ala Se 123 65 125 Arg Leu Ala Val Leu Eeu Ala Gly Thr Hi 126 85 127 61 100 127 100 131 Leu Gln Glu Leu Gly Ile Pro Gln Asg Leu 132 115 133 130 135 135 130 135 137 Gln Gly Ser Ser Leu Pro His Val Ser Ty 137 Gln Gly Ser Ser Leu Pro His Val Ser Ty	Ser Gln Pro Ser Pro Glu 30  Leu Asp Arg Ser Thr Phe 45  Leu His Gly Lys Asp Cys 60  r Ala Asn Leu Ser Glu Glu 75  80  s Thr Leu Leu Gln Gln Ala 95  O Asp Ala Phe Gln Glu Glu 110  u Ile Gly Asp Leu Ala Ser 125  u Asp Ser Val Ala Gln Gln 140  or Phe Arg Trp Arg Val Asp 160
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DATE: 07/26/2001 TIME: 18:43:12

#### Input Set : A:\1272591.app Output Set: N:\CRF3\07262001\1904568.raw

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 170 ggtttccccg cogcccacc ggacgccgac gaaagccagc gagctcctca gcctcaggca 240
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 176 tot got gtg ggg got goa act coa tac ctg cat cat cct ggt gat agt
 177 Ser Ala Val Gly Ala Ala Thr Pro Tyr Leu His His Pro Gly Asp Ser
 180 cac agt ggc cga gtg agt ttc ttg ggg gcc cag ctt cct cca gag gtg
  181 His Ser Gly Arg Val Ser Phe Leu Gly Ala Gln Leu Pro Pro Glu Val
  184 gca gca atg gcc cgg cta cta ggg gac cta gac agg agc acg ttc aga
                                                                         441
  185 Ala Ala Met Ala Arg Leu Leu Gly Asp Leu Asp Arg Ser Thr Phe Arg
  188 aag ttg ctg aag ttt gtg gtc agc agc ctg cag ggg gag gac tgc cga
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  189 Lys Leu Leu Lys Phe Val Val Ser Ser Leu Gln Gly Glu Asp Cys Arg
  192 gac ggt gtg cag cgt ctt ggg gtc agc gcc aac ctg ccg gag gag cag
                                                                         537
  193 Asp Gly Val Gln Arg Leu Gly Val Ser Ala Asn Leu Pro Glu Glu Gln
   196 ctg ggt gcc ctg ctg gca ggc atg cac aca ctg ctc cag cag gcc ctc
   197 Leu Gly Ala Leu Leu Ala Gly Met His Thr Leu Leu Gln Gln Ala Leu
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   201 Arg Leu Pro Pro Thr Ser Leu Lys Pro Asp Thr Phe Arg Asp Gln Leu
   204 cag gag ctc tgc atc ccc caa gac ctg gtc ggg gac ttg gcc agc gtg
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   208 gta ttt ggg agc cag cgg ccc ctc ctt gat tct gtg gcc cag cag cag
    209 Val Phe Gly Ser Gln Arg Pro Leu Leu Asp Ser Val Ala Gln Gln Gln
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# Input Set : A:\1272591.app Output Set: N:\CRF3\07262001\1904568.raw

Output Set: N:\CRF3\07262001\1904300:22	
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226 gag atg gca gat ctg gag aag agg tgt gag cgc agu atg 228 gag atg gca gat ctg gag aag acg cgc agu Arg Arg Leu Gln A: 229 Glu Met Ala Asp Leu Glu Lys Arg Cys Glu Arg Arg Leu Gln A: 220 220 220 220 220 220 220 220 220 220	spe
229 Glu Met Ala Asp bet 32 220 220 220 230 210 215 225 225 225 232 240 240 240 240 240 240 240 240 240 24	gatggtgc 1026
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253 Met Ser Ala Vul 57 10 10 254 1 5 5 30 256 Ser His Ser Gly Arg Val Ser Phe Leu Gly Ala Gln Leu Pro 25 25 25 25 25 25 25 25 25 25 25 25 25	110 014
256 Ser HIS Sel Call 25	Thr Phe
256 Ser H3 502 257 20 257 20 257 20 259 Val Ala Ala Met Ala Arg Leu Leu Gly Asp Leu Asp Arg Ser 45 40 45	
259 Val Ala State	Asp Cys
262 Arg Lys Leu Leu Lys Phe Val Val Ser 60	-1 (1)
262 Arg Lys Leu Sco 275 55 60 263 50 263 50 265 Arg Asp Gly Val Gln Arg Leu Gly Val Ser Ala Asn Leu Pro 75 70 70 70 70 70 70 70 70 70 70 70 70 70	80
265 Arg Asp Gly Val Gln 70	Gln Ala
265 Arg Asp Gly Val San 70 266 65 70 268 Gln Leu Gly Ala Leu Leu Ala Gly Met His Thr Leu Leu Gln 90 80 Rep Dec Arg	95
268 GIn Leu Gly All 85 90 90 269 85 271 Leu Arg Leu Pro Pro Thr Ser Leu Lys Pro Asp Thr Phe Arg 271 Leu Arg Leu Pro Pro Thr Ser Leu Lys Pro Asp 100	Asp Gln
271 Leu Arg Leu Pro Pro Thr Ser Leu Lys Plo ASP 1110	
271 Leu Gln Glu Leu Cys Ile Pro Gln Asp Leu Val Gly Asp Leu 272 100 1120 120 125	Ala Ser
274 Leu Gln Glu Leu Cys IIe Plo Gli May	ain Cin
275 115 Cor Cin Arg Pro Leu Leu Asp Ser Val Ala	GIN GIN
274 Leu Gir Gia Garage 120 275 115 120 277 Val Val Phe Gly Ser Gin Arg Pro Leu Leu Asp Ser Val Ala 277 Val Val Phe Gly Ser Gin Arg 135	Val ASP
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280 Gin Gly All 150 150 150 200 Len Gln Pro	Ser Val
280 GH GJ Mad 31 150 150 281 145 283 Val Ala Ile Ser Thr Ser Ala Leu Ala Arg Ser Leu Gln Pro	175
284 165	

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293		210					213									

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Input Set : A:\1272591.app Output Set: N:\CRF3\07262001\1904568.raw

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